This listing of claims will replace all prior versions and listings of claims in the application:

Listing of Claims:

- (currently amended) An isolated polypeptide comprising a <u>BST Over-expressed</u> Gene (BOG) BOG polypeptide-fragment, said BOG <u>polypeptide-fragment</u> comprising
 - (i) at least about 90% amino acid sequence identity with SEQ ID NO: 8;
 - (ii) a retinoblastoma gene product (pRB) a pRb binding motif; and
 - (iii) at least one a-casein kinase II phosphorylation motif;

wherein the polypeptide binds pRB and displaces E2F-1 bound to pRB.

2-3. (cancelled)

- (currently amended) The BOG polypeptide fragment of claim 1, wherein said casein kinase II phosphorylation motif is located downstream of the pRb binding motif.
- (currently amended) The BOG polypeptide fragment of claim 4, further comprising a second casein kinase II phosphorylation motif, said second casein kinase II phosphorylation motif being located upstream of the pRb binding motif.
 - (currently amended) The BOG polypeptide fragment of claim 1 joined to a
 detectable label.
- (currently amended) The BOG polypeptide fragment of claim 6, wherein the detectable label includes a radioactive isotope, an enzyme, a chromophore or a mixture thereof.

8-22. (cancelled)

 (currently amended) A chimeric molecule comprising a BOG polypeptide of claim 1 fragment fused to a heterologous amino acid sequence.

24-32. (cancelled)

iii)

- 33. (currently amended) An isolated polypeptide comprising:
- i) at least 90% amino acid sequence identity with SEQ ID NO:2, SEQ ID NO:8 or SEO ID NO:10:
- ii) a retinoblastoma gene product produce (pRB) binding motif; and at least one casein kinase II phosphorylation motif:
- wherein the polypeptide binds pRB and displaces E2F-1 bound to pRB.
- 34. (currently amended) The polypeptide of claim 33 comprising the an-amino acid sequence of SEQ ID NO:2.
- 35. (currently amended) The polypeptide of claim 33 comprising the an-amino acid sequence of SEO ID NO:8.
- 36. (currently amended) The polypeptide of claim 33 comprising the an-amino acid sequence of SEO ID NO:10.
- 37. (previously presented) The polypeptide of claim 33, wherein said casein kinase II phosphorylation motif is located downstream of the pRB binding motif.
- 38 (previously presented) The polypeptide of claim 37 further comprising a second casein II phosphorylation motif, said second casein kinase II phosphorylation motif being located upstream of the pRB binding motif.
 - 39. (previously presented) The polypeptide of claim 33 joined to a detectable label.
- 40 (previously presented) The polypeptide of claim 39, wherein the detectable label comprises a radioactive isotope, an enzyme, a chromophore or a mixture thereof.

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- 41. (previously presented) The polypeptide of claim 33 further comprising a heterologous amino acid sequence.
- 42. (previously presented) The polypeptide of claim 41, wherein the heterologous amino acid is a tag polypeptide.
- (previously presented) The polypeptide of claim 41, wherein the heterologous amino acid sequence is that of an immunoglobulin constant region.
- (previously presented) The polypeptide of claim 41, wherein the heterologous amino acid sequence is maltose binding protein.
- 45. (new) An isolated polypeptide comprising a polypeptide having 95% sequence identity to the amino acid sequence of SEQ ID NO:8, wherein the isolated polypeptide binds pRB and displaces E2F-1 bound to pRB.
- (new) The isolated polypeptide of claim 45, comprising the amino acid sequence of SEQ ID NO:8.
- 47. (new) The isolated polypeptide of claim 1, comprising a polypeptide having 95% sequence identity to the amino acid sequence of SEQ ID NO:8.
- 48. (new) The isolated polypeptide of claim 47, comprising the amino acid sequence of SEO ID NO:8.